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The new GeneChip\* One-Cycle and Two-Cycle cDNA Synthesis Kits.



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# **# GETTING** STARTED

-> Wizard

### :: QUERY Expression

- -> Quick Query
- -> Standard Query
- -> Batch Query
- -> BLAST
- -> Probe Match
- -> UCSC Query

## Genotyping

- -> Quick Query
- -> Standard Query
- -> Batch Query
- -> UCSC Query
- -> SNP Finder

## **# CURRENT QUERY** 1 probe sets

- -> Annotations
- -> Show Orthologs
- -> GO Browser
- -> Export

## **# QUERY HISTORY**

#### **Annotation Views**

- -> Expression
- -> Genotyping

# -> BLAST Status

# -> New Folder -> Expression

- Queries -> (1)All Descriptions
- (AFFX-BioDn-5 st) -) (1) All Descriptions
- (AFFX-BioB-M st) → (1)All Descriptions
- (HG613) -) (2)All Descriptions (AFFX-BioDn-5)
- (2)All Descriptions (AFFX-BioB-M)
- -> Genotyping Queries

## **Full Record**

# Details for HUGENEFL: AFFX-BIODN-5\_ST

**Full Screen** 

NetAffx Links

Cluster Members Consensus/Exemplar

GeneChip Array Information

Probe Set ID AFFX-BioDn-5\_st

GeneChip HumanGeneFL Array Array

Organism

Common Human

Name

Target

Sequence

# **Probe Design Information**

Transcript ID AFFX-BioDn-5

Sequence Control sequence Type

Representative J04423 NCBI Public ID

J04423 E coli bioD gene dethiobiotin synthetase (-5 and -3 represent transcript Target regions 5 prime and 3 prime respectively) Description

# Sequence

>HUGENEFL:AFFX-BIODN-5 ST

gggaaaactgtcgccagttgtgcacttttacaagccgcaaaggcagcaggctaccggacg gcaggttataaaccggtcgcctctggcagcgaaaagaccccggaaggtttacgcaatagc  $\tt gacgcgctggcgttacagcgcaacagcagcctgcagctggattacgcaacagtaaatcct$ tacaccttcgcagaacccacttcgccgcacatcatcagcgcgcaagagggcagaccgata

gaatcattggtaatgagcgccggattacgcgcgcttg

	Probe Sequence(5'-3')	Probe X	Probe Y	Probe Interrogation Position	Strandedness
	GTGCACAACTGGCGACAGTTTTCCC	281	11	49	Sense
	GGCTTGTAAAAGTGCACAACTGGCG	282	11	60	Sense
	GCTGCCTTTGCGGCTTGTAAAAGTG	283	11	71	Sense
	GGTAGCCTGCTGCCTTTGCGGCTTG	284	11	79	Sense
Probe Info	CCGTCCGGTAGCCTGCTGCCTTTGC	285	11	85	Sense
	CAGCGCGTCGCTATTGCGTAAACCT	286	11	153	Sense
	GTAACGCCAGCGCGTCGCTATTGCG	287	11	160	Sense
	TTGCGCTGTAACGCCAGCGCGTCGC	288	11	167	Sense
	TGCTGTTGCGCTGTAACGCCAGCGC	289	11	172 ·	Sense
	TGCAGGCTGCTGTTGCGCTGTAACG	290	11	179	Sense

TCCAGCTGCAGGCTGCTGTTGCGCT	291	11	185	Sense
TGCGTAATCCAGCTGCAGGCTGCTG	292	11	192	Sense
TTACTGTTGCGTAATCCAGCTGCAG	293	11	199	Sense
CGGTCTGCCCTCTTGCGCGCTGATG	294	11	261	Sense
GATTCTATCGGTCTGCCCTCTTGCG	295	11	269	Sense
TACCAATGATTCTATCGGTCTGCCC	296	11	276	Sense
CTCATTACCAATGATTCTATCGGTC	297	11	281	Sense
TCCGGCGCTCATTACCAATGATTCT	298	11	288	Sense
CGCGTAATCCGGCGCTCATTACCAA	299	11	295	Sense
CAAGCGCGCGTAATCCGGCGCTCAT	300	11	301	Sense

888-DNA-CHIP (888-362-2447) | +44 (0) 1628 552550

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